

FIG. 1A-1
FIG. 1A-2

FIG. 1A

GAATTCCCCAACAGAGCCAAGCTCTCCATCTAGTGGACAGGGAAGCTAGCAGCAACC	39 (UPPER: SEQ ID NO.: 1)
	19 (LOWER: SEQ ID NO.: 4)
TTCCCTTCACCTACAAACTTCATTGCTTGGCCAAAAGAGAGTTAATTCATGTAGACAT	119
	39
CTATGTAGGCAATTAAAAACCTATTGATGTATAAAACAGTTTGCAATTCATGGAGGGCAAC	179
	59
TAAATACATTCTAGGACTTTATAAAGATCACTTTTATTATGTCACAGGGTGGAAACAAG	239
	79
ATGGATTATCAAGTGTCAAGTCCAAATCTATGACATCAATTAATTATACATCGGAGCCCCTGC	299
M D Y Q V S S P I Y D I N Y Y T S E P C	99

1/20

FIG. 1A-1

CAAAAAATCAATGTGAAGCAAAATCGCAGCCCGCCTCCTGCCCTCCGCTCTACTCACTGGTG	359
Q K I N V K Q I A A R L L P P L Y S L V	119
TTCATCTTTGGTTTGTGGGCAACATGCTGGTCATCCTCATCCTGATAAACTGCAAAAGG	419
F I F G F V G N M L V I L I L I N C K R	139
CTGAAGAGCATGACTGACATCTACCTGCTCAACCTGGCCATCTCTGACCTGTTTTTCCTT	479
L K S M T D I Y L L N L A I S D L F F L	159
CTTACTGTCCCTTCTGGGCTCACTATGCTGCCGCCAGTGGGACITTTGGAATACAATG	539
L T V P F W A H Y A A A Q W D F G N T M	179
TGTCAACTCTTGACAGGGCTCTATTTTATAGGCTTCTTCTCTGGAATCTTCTTCATCATC	599
C Q L L T G L Y F I G F F S G I F F I I	199
CTCCTGACAATCGATAGGTACCTGGCTGTCGTCATGCTGTGTTTTCCTTAAAGCCAGG	659
L L T I D R Y L A V V H A V F A L K A R	219
ACGGTCACCTTTGGGTTGTGACAAGTGTGATCACTTGGGTGGTGGCTGTGTTGCGTCT	719
T V T F G V V T S V I T W V V A V F A S	239
CTCCAGGAATCATCTTTACCAGATCTCAAAAAGAGTCTTTCATTACACCTGCAGCTCT	779
L P G I I F T R S Q K E G L H Y T C S S	259
CATTTTCCATACA	
H F P Y	

FIG. 1A-2

FIG. 1B-1
FIG. 1B-2

FIG. 1B

GAATTCCCCCAACAGAGCCAGCTCTCCATCTAGTGGACAGGGAAGCTAGCAGCAAACC	59 (UPPER: SEQ ID NO.: 2)
	19 (LOWER: SEQ ID NO.: 5)
TTCCCTTCACACAAAACCTTCATTGCTTGSCCAAAAGAGAGTTAATTCAATGTAGACAT	119
	39
CTATGTAGGCAATTAAAAACCTATTGATGTATAAAACAGTTTGCATTTCATGGAGGGCAAC	179
	59
TAAATACATTCTAGGACCTTTATAAAAGATCACCTTTTATTATGACAGGGTGGAACAAG	239
	79
ATGGATTATCAAGTGTCAAGTCCAATCTATGACATCAATTATTATATACATCGGAGCCCTGC	299
M D Y Q V S S P I Y D I N Y Y T S E P C	99

FIG. 1B-1

CAAAAATCAATGTGAAGCAAAATCGAGCCCGCCTCCTGCCTCCGCTCTACTCACTGGTG 359
Q K I N V K Q I A A R L L P P L Y S L V 119

TTCACTCTTTGGTTTGTGGGCAACATGCTGGTCATCCTCATCCTGATAAACTGCAAAAGG 419
F I F G F V G N M L V I L I L I N C K R 139

CTGAAGAGCATGACTGACATCTACTCTGCTCAACCTGGCCATCTCTGACCTGTTTTCCTT 479
L K S M T D I Y L L N L A I S D L F L 159

CTTACTGTCCCTTCTGGGCTCACTATGCTGCGGCCAGTGGGACTTTGGAAATACAATG 539
L T V P F W A H Y A A A Q W D F G N T M 179

TGTCAACTCTTGACAGGGCTCTATTTTATAGGCTTCTCTCTGGAATCTTCTTTCATCATC 599
C Q L L T G L Y F I G F F S G I F I I 199

CTCCTGACAAATCGATAGGTACCTGGCTGTGTCCTCATGCTGTGTTTGTAAAGCCAGG 659
L L T I D R Y L A V V H A V F A L K A R 219

ACGGTCACCTTTGGGGTGGTGACAAGTGTGATCACTTGGGTGGTGGCTGTGTTTGCCTCT 719
T V T F G V V T S V I T W V V A V F A S 239

CTCCCAGGAATCATCTTTACCAGATCTCAAAAAGAAAGTCTTCAATTACACCTGCAGCTCT 779
L P G I I F T R S Q K E G L H Y T C S S 259

CATTTTCCATACAGTCAGTATCAATTTCTGGAAGAAATTTCCAGACATTTAAAGATAGTCATC 839
H F P Y S Q Y Q F W K N F Q T L K I V I 279

FIG. 1B-2

TTGGGGCTGGTCTCTGCCGCTGCTTGTGTCATGGTCACTGCTACTCGGGAATCCTAAAAACT 899
 L G L V L P L L V M V I C Y S G I L K T 299
 CTGCTTCGGTGTGCGAAATGAGAAGAGAGGCACAGGGCTGTGAGGCTTATCTTCACCATC 959
 L L R C R N E K K R H R A V R L I F T I 319
 ATGATTGTTTATTTTCTCTTCTGGGCTCCCTACAACATTGTCTCTTCTCTGAAACACCTTC 1019
 M I V Y Y F L F W A P Y N I V L L L N T F 339
 CAGGAATCTTTGGCCTGAATAATTGCAGTAGCTCTAACAGGTTGGACCAAGCTATGCAG 1079
 Q E F F G L N N C S S S N R L D Q A M Q 359
 GTGACAGAGACTCTTTGGGATGACGCACCTGTGTCATCAACCCCATCATCTATGCCCTTGTGTC 1139
 V T E T L G M T H C C I N P I I Y A F V 379
 GGGGAGAAGTTCAGAAACTACCTCTTAGTCTCTTCTTCCAAAGCACATTGCCAAACGCTTC 1199
 G E K F R N Y L L V F F Q K H I A K R F 399
 TGCAAAATGCTGTTCTATTTTCCAGCAAGAGGCTCCCGAGCGAGCAAGCTCAGTTTACACC 1259
 C K C C S I F Q Q E A P E R A S S V Y T 419
 CGATCCACTGGGGAGCAGGAAATATCTGTGGGCTTGTGACACGGACTCAAGTGGGCTGGT 1319
 R S T G E Q E I S V G L * 439
 GACCCAGTCAGAGTTGTGCACATGGGCTTAGTTTTCATACACAGCCTGGGCTGGGGTNGG 1379
 459
 TTGGNNGAGGTCTTTTTTAAAGGAAGTTACTGTATTAGAGGGTCTAAGATTTCATCCATT 1439
 479
 TATTTGGCATCTGTTTAAAGTAGATTAGATCCGAATTC

FIG. 1B-3

FIG. 1D-1
FIG. 1D-2

FIG. 1D

GAATTCCCCAACAGAGCCCAAGCTCTCCATCTAGTGGACAGGGAAGCTAGCAGCAACC	59 (UPPER: SEQ ID NO. 3) 19 (LOWER: SEQ ID NO. 6)
TTCCCTTCACTACAAAACCTTCATTGCTTGGCCAAAAGAGAGGTTAAATTCATGTAGACAT	119 39
CTATGTAGGCAATTAAAAACCTATTGATGTATAAACAGTTTGCAATTCATGGAGGGCAAC	179 59
TAAATACATTCTAGGACTTTATAAAAGATCACTTTTATTATGCACAGGGTGGAAACAAG	239 79
ATGGATTATCAAGTGTCAAGTCCCAATCTATGACATCAATATTATATACATCGGAGCCCTGC	299 99
M D Y Q V S S S P I Y D I N Y Y T S E P C	

FIG. 1D-1

CAAAAATCAATGTGAAGCAATCGCAGCCGCTCTGCTCCGCTCTACTCACTGTG 359
 Q K I N V K Q I A A R L L P P L Y S L V 119
 TTCATCTTTGGTTTGTGGCAACATGCTGGTCATCCTCATCAATAAAGG 419
 F I F G F V G N M L V I L I L I N C K R 139
 CTGAAGAGCATGACTGACATCTACCTGCTCAACCTGGCCATCTCTGACCTGTTTCCCT 479
 L K S M T D I Y L L N L A I S D L F F L 159
 CTTACTGTCCCTTCTGGGCTCACTATGCTGCCGCCAGTGGGACTTTGGAATACAATG 539
 L T V P F W A H Y A A A Q W D F G N T M 179
 TGTCAACTCTTGACAGGGCTCTATTTTATAGGCTTCTTCTGGAATCTTCTTCATCATC 599
 C Q L L T G L Y F I G F F S G I F F I I 199
 CTCCTGACAAATCGATAGGTACCTGGCTGCTGCTCCATGCTGTGTTTAAAGCCAGG 659
 L L T I D R Y L A V V H A V F A L K A R 219
 ACGGTCACTTTGGGTGGTGACAAGTGTGATCACTTGGGTGGTGGTGTGTTGCGTCT 719
 T V T F G V V T S V I T W V V A V F A S 239
 CTCCCAGGAATCATCTTTACCAGATCTCAAAAGAAGGTCTTCAATTACACCTGCAGCTCT 779
 L P G I I F T R S Q K E G L H Y T C S S 259
 CATTTCCATACATTAAAGATAGTCATCTTGGGGTGGTCCCTGCCGTGCTTGTGATGTT 839
 H F P Y I K D S H L G A G P A A A C H G 279

FIG. 1D-2

CATCTGCTACTCGGGAATCCTAAAACTCTGCTTCGGTGTGAAATGAGAAGAGGCA	899
H L L L G N P K N S A S V S K *	299
CAGGGCTGTGAGGCTTATCTTCACCATCATGATTGTTTATTTCTCTCTCGGGCTCCCTA	959
	319
CAACATGTCTCTTCTCTGAACACCTTCCAGGAATTCCTTGGCCTGAATAATTCAGTAG	1019
	339
CTCTAACAGGTTGGACCAAGCTATGCAGGTGCAGAGACTCTTGGGATGACGCACCTGCTG	1079
	359
CATCAACCCCATCATCTATGCCTTTGTGCGGGGAGAAGTTCAGAACTACCTCTTAGTCTT	1139
	379
CTTCCAAAAGCACATTGCCAAACGCTTCTGCAAATGCTGTTCTATTTCACGCAAGAGGC	1199
	399
TCCCGAGCGAGCAAGCTCAGTTTACACCCGATCCACTGGGGAGCAGGAAATATCTGTGGG	1259
	419
CTTGTGACACGGACTCAAGTGGGCTGGTGACCCAGTCAGAGTTGTGCACATGGCTTAGTT	1319
	439
TTCATACACAGCCTGGGCTGGGGTNGGTTGNNAGGCTTTTTTAAAGGAAGTTACT	1379
	459
GTTATAGAGGGTCTAAGATTCTATCCATTTATTTGGCATCTGTTTAAAGTAGATTAGATCC	1439
	479
GAATTC	

FIG. 1D-3

FIG. 2A
FIG. 2B

FIG. 2

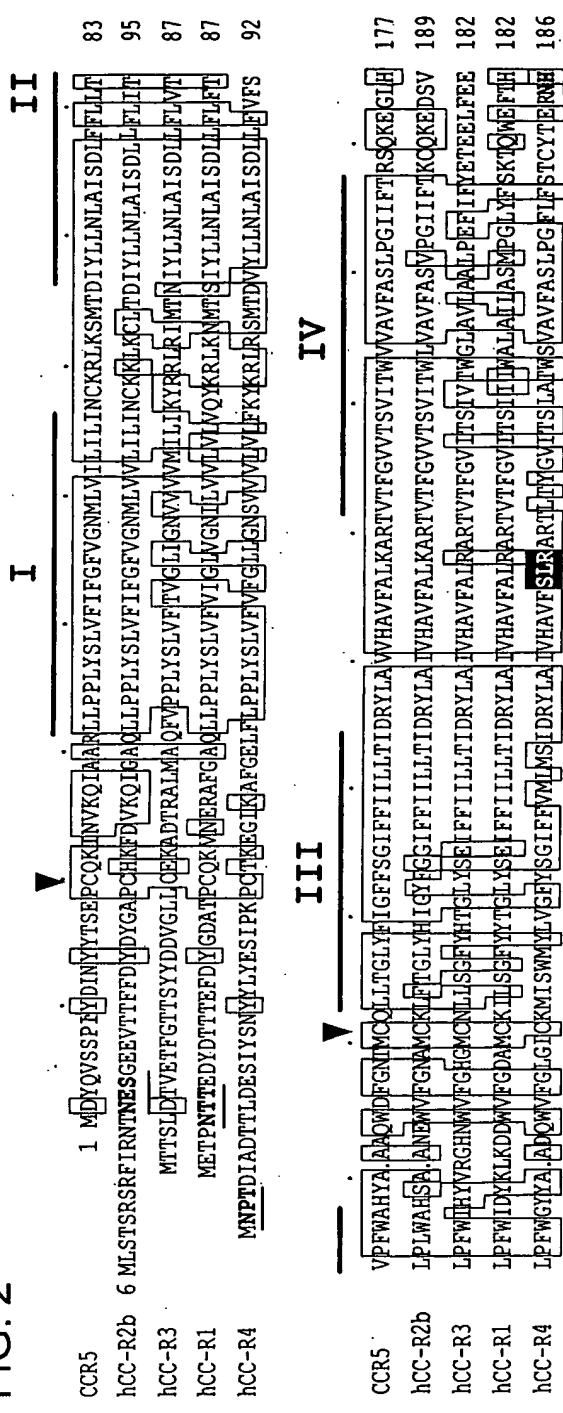


FIG. 2A

VI

V

CCR5 YTCSEHFPYSQYQFWKNEQTLKIYILGLVPLLVVVICYSGLKTLRCRNEKKRRRAVRLIFTIMIVYFLFWAPYNNIVLLNLTQEFFGLNNC 272
 hcc-R2b YVQGPYFPRG...WNNFHTIMRNILGLVPLLVVICYSGLKTLRCRNEKKRRRAVRIFTIMIVYFLFWAPYNNIVLLNLTQEFFGLSNC 280
 hcc-R3 TICSALYPEDTVYSWRHHHTLRMTIFQLVPLLVVNAICVYTGHTKTLRCPSKKK.YKAIQRLLIFVIMAVFFFIFFWTPYNNVAILLSSVQSILFGNDQ 276
 hcc-R1 HTCSHFFPHESLREWKLPQALKNLNLGLVPLLVVVICVYTGHTKTLRCRNEKK.SKAVRLIFVIMIHFFELFWTPYNNLTILISVFQDELFTHEC 276
 hcc-R4 TYCKTKYSINST.TWKVLSLSLEINILGLVPLLVVICYSGLKTLRCRNEKK.NKAVKMFPAVVVFLGFWTPYNNIVLFIETLVLELVLDQDC 279

VII

CCR5 SSSNRLDQAMQVTETLGMTHCCINPIIYAFVGEKFRNLLVFFQKHIAGR.FCKCCSIFQOEAFERASSVYTRNSTGEOEISVGL 352
 hcc-R2b ESTSQLDQAQVETETLGMTHCCINPIIYAFVGEKFRNLLVFFQKHIAGR.FCKQCPVEYRETVDGVTSNIPSTGEOEWSAGL 360
 hcc-R3 ERSKHLDLVMILVTEVIAYSHCCNPNIIYAFVGEKFRKVIHRHFRHLIMH.IGRYIPFIPSEKIERTSV.SPSTAFPELSIVF 355
 hcc-R1 EQSRHLDLVMQVTEVIAYTHCCVNPPIIYAFVGEKFRKVIHQLFHRRVAVH.LVMWLPFLSVDRUERVSST.SPSTGEHELSAGF 355
 hcc-R4 TFERYLDVYATATETIAFVHCCLNPIIYFELGKFRKVIHQLFHRRVAVH.LVMWLPFLSVDRUERVSST.SPSTGEHELSAGF 360

FIG. 2B

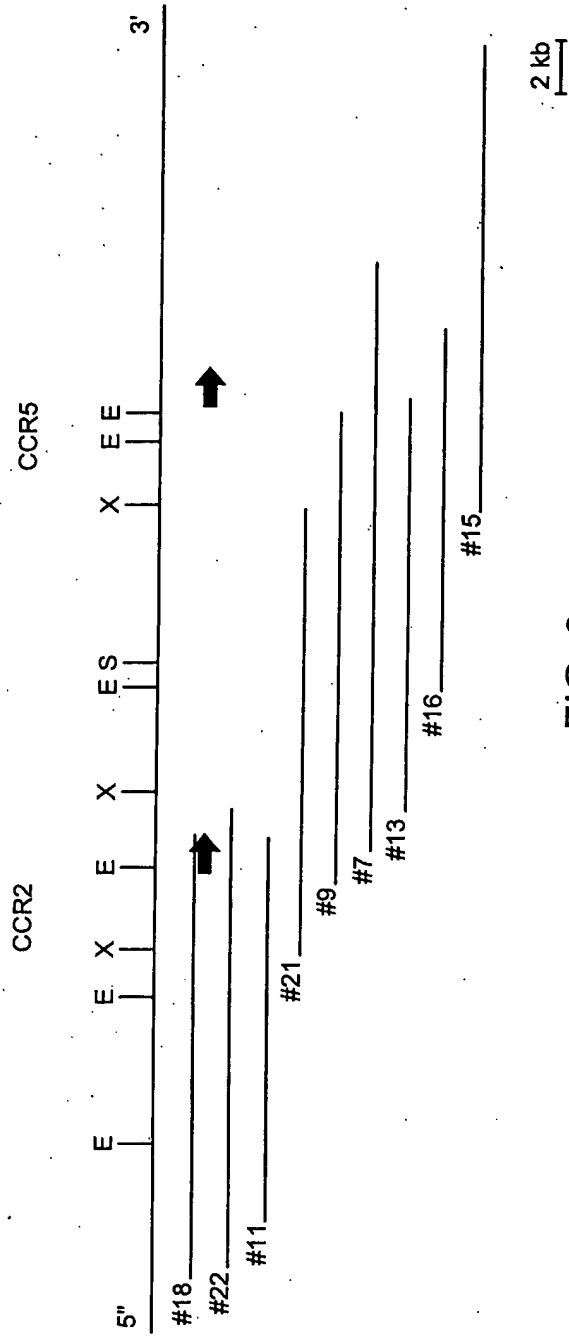


FIG. 3

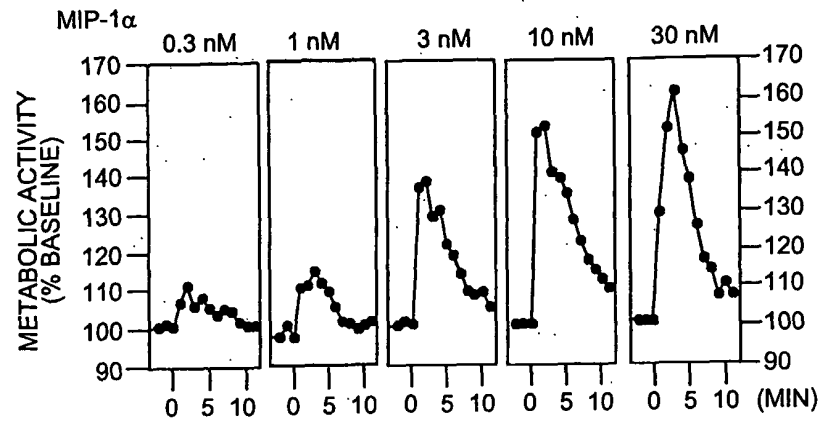


FIG. 4A

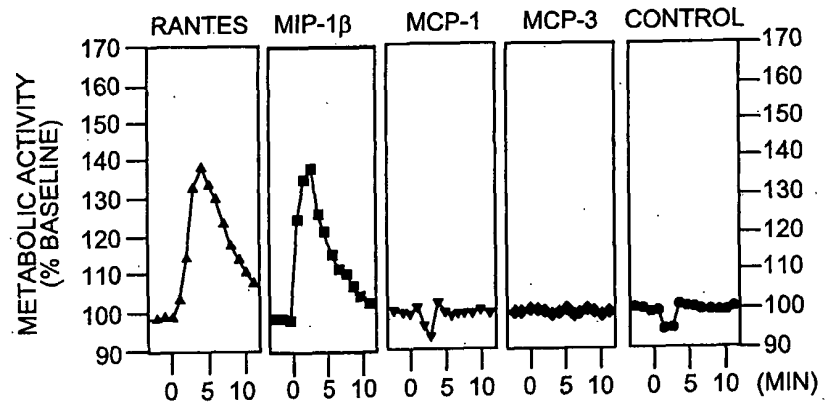


FIG. 4B

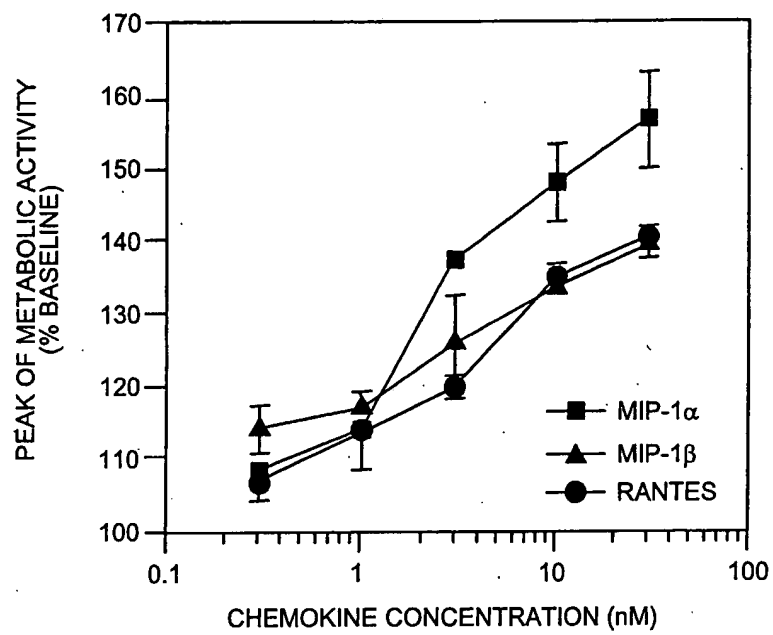


FIG. 4C

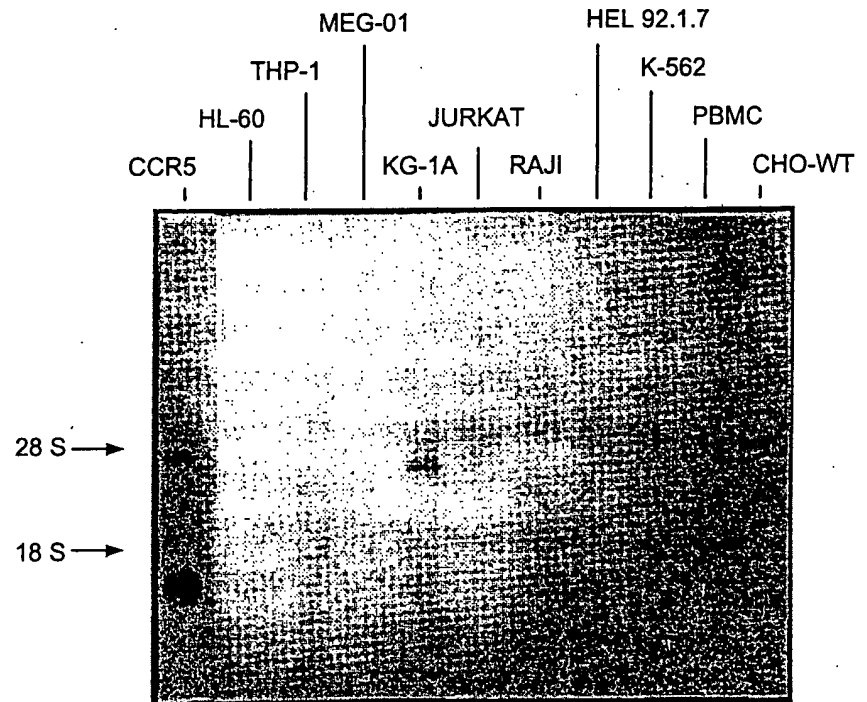
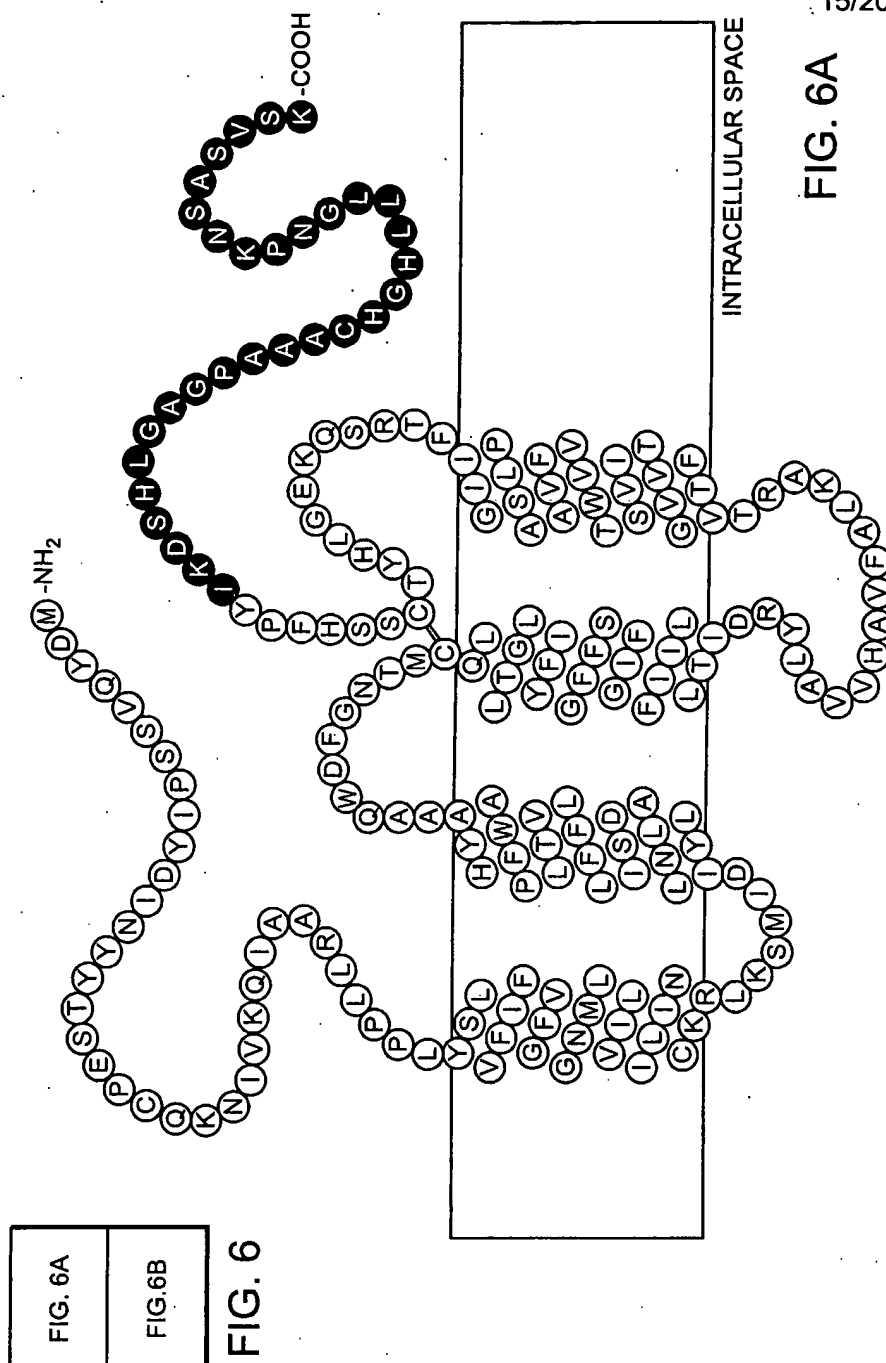


FIG. 5

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FIG. 6A




CCR5 F P Y S Q Y Q F W K N F Q T L K I V I L G L V L P
~~TTTCCATACagtcagtcataattctggaagaatttccagacaTTAAAGATAGTCATCTTGGGGTGGTCTCTGCGCG~~
 Δccr5 F P Y  I K D S H L G A G P A
 CCR5 L L V M V I C Y S G I L L K T L L R C R N E K K R
 CTGCTTGTCATGGTCATCTGCTACTCGGAATCCTAAAACCTGCTTCGGTTCGAATGAGAAGAAGG
 Δccr5 A A C H G H L L L L G N P K N S A S V S K *

FIG. 6B

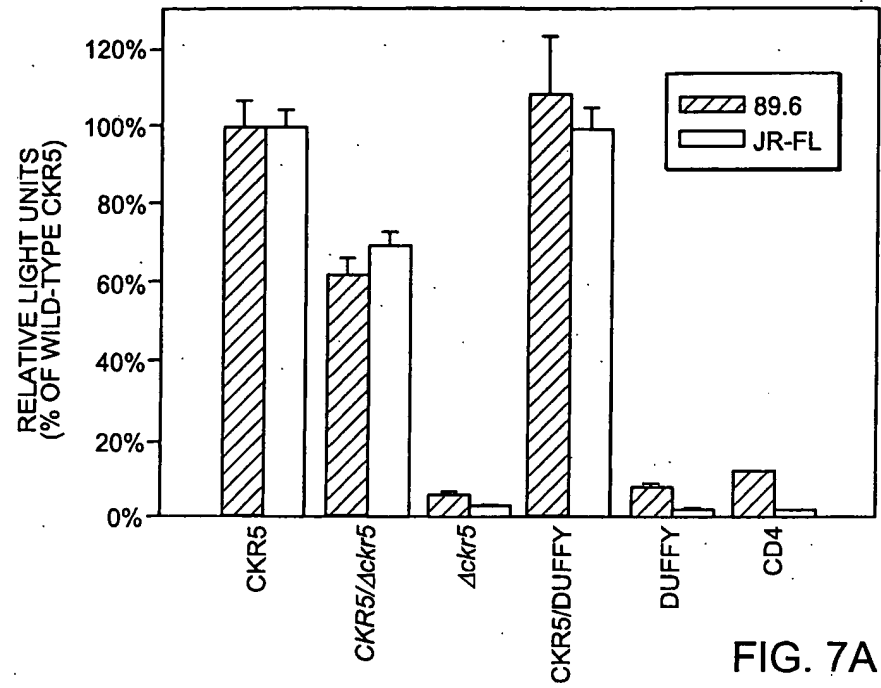


FIG. 7A

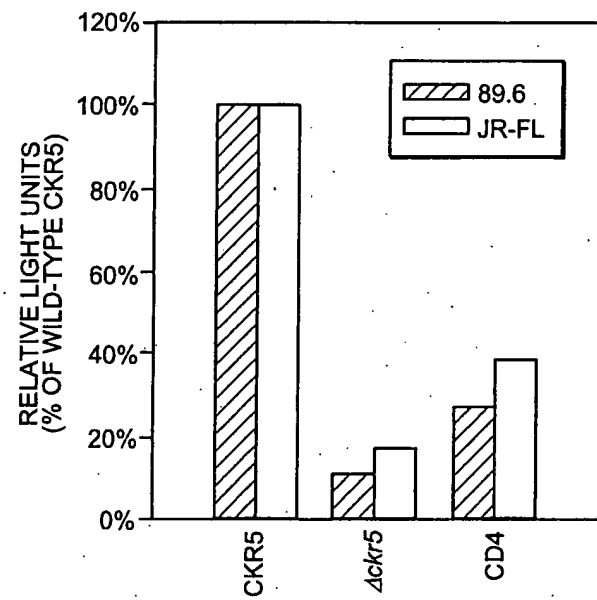


FIG. 7B

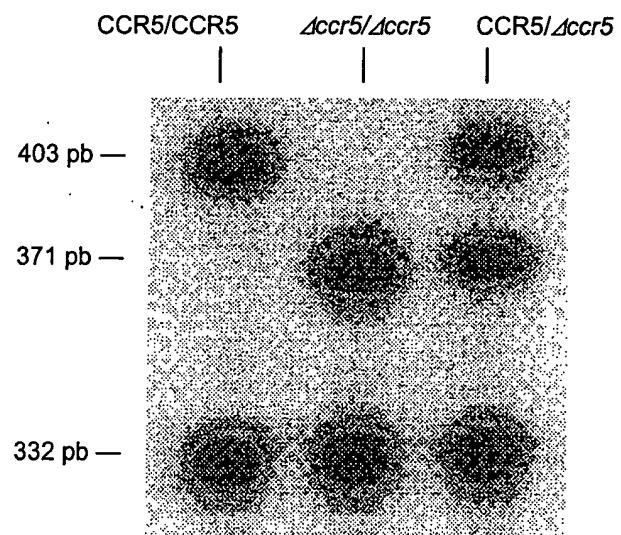
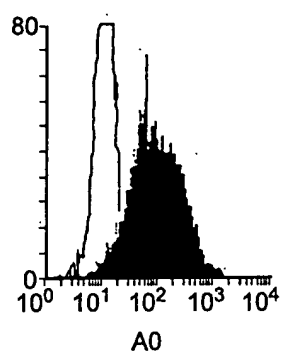
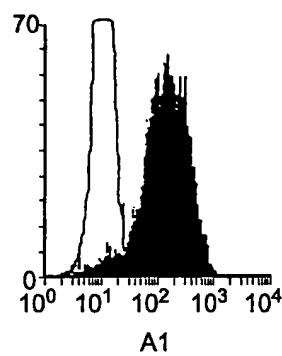
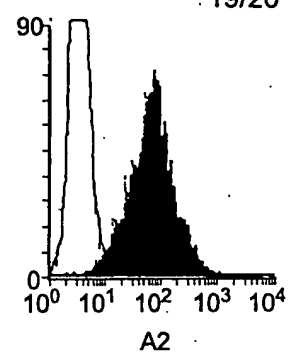
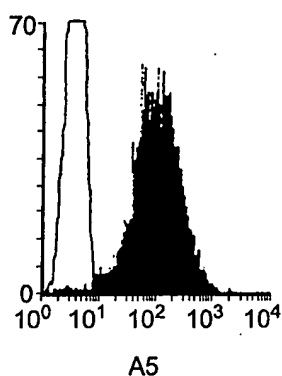
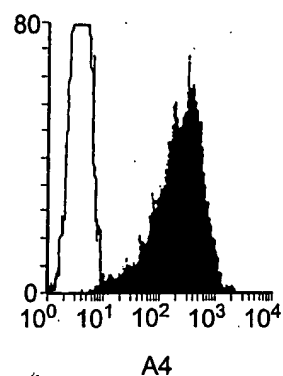
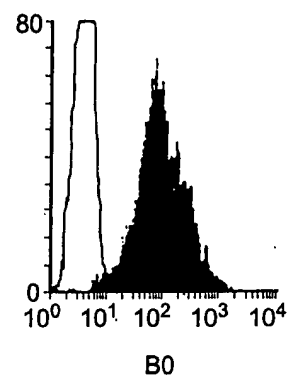
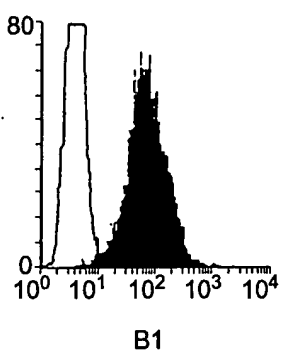
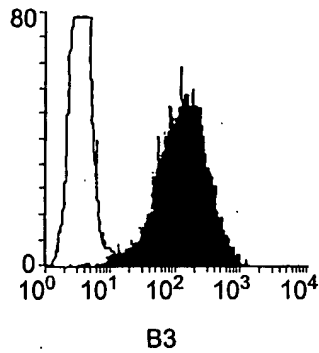


FIG. 8

A0
FIG. 9AA1
FIG. 9BA2
FIG. 9CA5
FIG. 9DA4
FIG. 9EB0
FIG. 9FB1
FIG. 9GB3
FIG. 9H

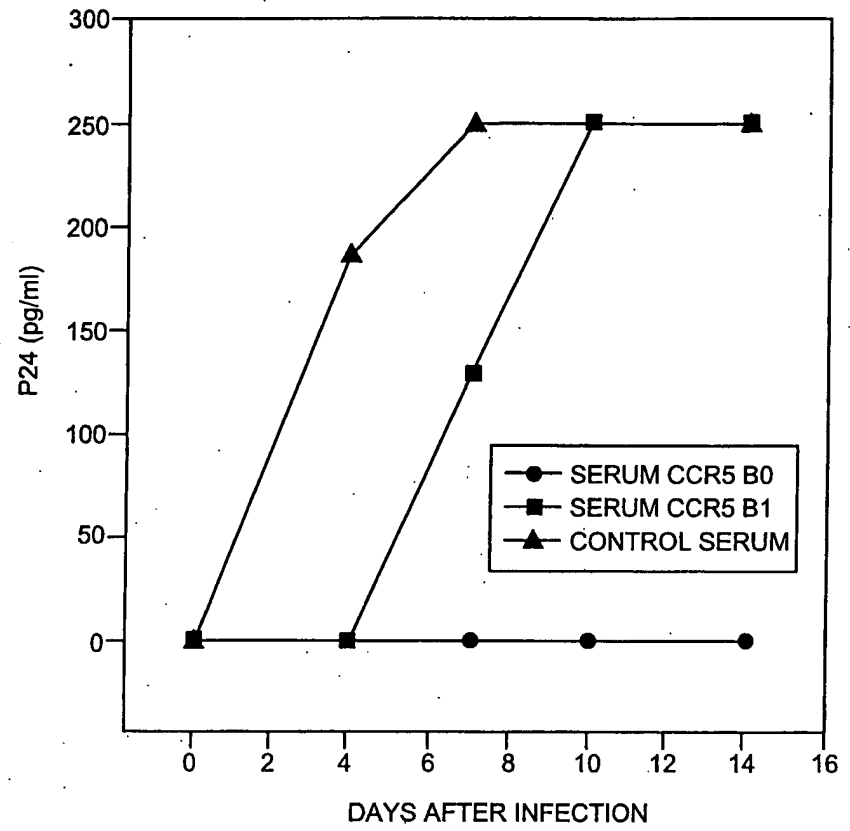


FIG. 10